NUCLEOTIDE AND AA SEQUENCE

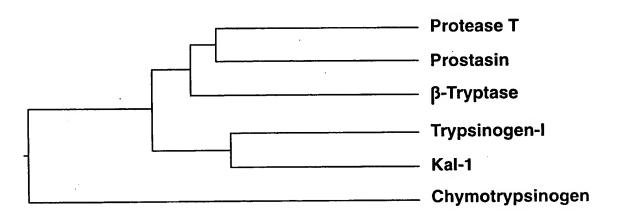
Nucleic acid sequence of Protease T (SEQ.ID.NO.:1)

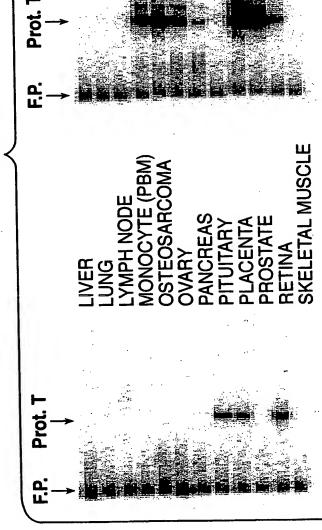
GACCACGCCCTGCGCCCAGCCAGGCCTGAGGACATGAGGCGGCCGCCGCGGGGGCGC TCCTGCTGCTGTGTTTTTGGGTCTCAGAGGCCAAGGCAGCAACAGCCTGTGGTCGCC CCAGGATGCTGAACCGAATGGTGGGCGGCCAGGACACGCAGGAGGGCGAGTGGCCCTGGC AAGTCAGCATCCAGCGCAACGGAAGCCACTTCTGCGGGGGCAGCCTCATCGCGGAGCAGT GGTCCTGACGGCTGCGCACTGCTTCCGCAACACCTCTGAGACGTCCCTGTACCAGGTCC AGGIGGAGAGCAACCCCTGTACCAGGGCACGCCTCCAGCGCTGACGIGGCCCTGGTGG AAGACCTCCTGCCGAACCGCGGATCCTGCAGAAACTCGCTGTGCCCATCATCGACACAC CCAAGTGCAACCTGCTCTACAGCAAAGACACCGAGTTTGGCTACCAACCCAAAACCATCA AGAATGACATGCTGTGCGCCGGCTTCGAGGAGGGCCAAGAAGGATGCCTGCAAGGGCGACT CGGCGCCCCTGGTGTGCCTCGTGGTCAGTCGTGGCTGCAGGCGGGGGTGATCAGCT GGGTGAGGGCTGTGCCCGCCAGAACCGCCCAGGTGTCTACATCCGTGTCACCGCCCACC AGAAGTGAGACCCCCGGGGCCAGGAGCCCCTTGAGCAGAGCTCTGCACCCAGCCTGCCCG CCCACACCATCCTGCTGTCCTCCCAGCGCTGCTGTTGCACCTGTGAGCCCCACCAGACT CCAATAAAAACCCAGCCTGTGTGCCAGCTG

>Amino Acid Sequence of Protease T (SEQ.ID.NO.:7)

MRRPAAVPLLLLLCFGSQRAKAATACGRPRMLNRMVGGQDTQEGEWPWQVSIQRNGSHFC GGSLIAEQWVLTAAHCFRNTSETSLYQVLLGARQLVQPGPHAMYARVRQVESNPLYQGTA SSADVALVELEAPVPFTNYILPVCLPDPSVIFETGMNCWVTGWGSPSEEDLLPEPRILQK LAVPIIDTPKCNLLYSKDTEFGYQPKTIKNDMLCAGFEEGKKDACKGDSGGPLVCLVGQS WLOAGVISWGEGCARQNRPGVYIRVTAHHNWIHRIIPKLQFQPARLGGQK

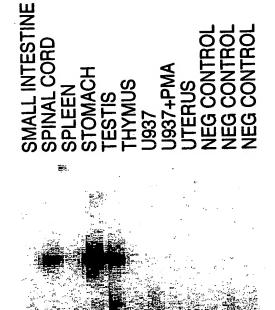
FIG. 2

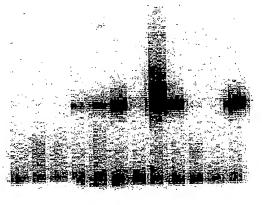




BCELL LYMPHOMA BONE MARROW AMYGDALA CAUDATE NUCLEUS CEREBELLUM CEREBELLUM CEREBRAL CORTEX HIPPOCAMPUS HYPOTHALAMUS SUBSTANTIA NIGRA SUBSTANTIA NIGRA SUBTHALMIC NUCLEUS THALAMUS BREAST CARCINOMA

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COLON ENDOTHELIAL EPIDERMIS FAT CELL FETAL BRAIN FIBROBLAST HEART JURKAT JURKAT KERATINOCYTE KIDNEY

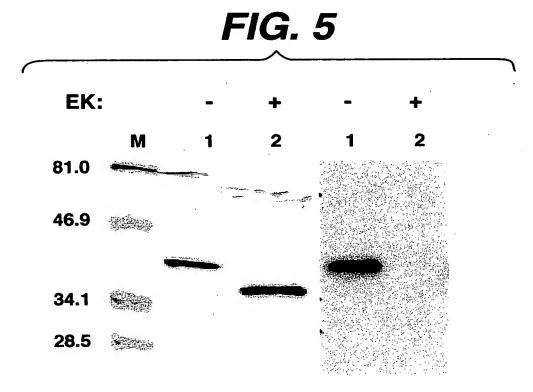
FUSION PROTEIN

Nucleic Acid sequence of PFEK-PROTT-HIS fusion protein (SEQ.ID.NO.:8).

GAATTCACCACCATGGACAGCAAAGGTTCGTCGCAGAAATCCCGCCTGCTCCTGCTGCTG GTGGTGTCAAATCTACTCTTGTGCCAGGGTGTGGTCTCCGACTACAAGGACGACGACGAC GTGGACGCGCCGCTCTTGCTGCCCCCTTTGATGATGATGACAAGATCGTTGGGGGCTAT CTCTAGAGGAGGCGAGTGGCCTGGCAAGTCAGCATCCAGCGCAACGGAAGCCACTTC TGCGGGGCAGCCTCATCGCGGAGCAGTGGGTCCTGACGGCTGCGCACTGCTTCCGCAAC ACCTCTGAGACGTCCCTGTACCAGGTCCTGCTGGGGGCAAGGCAGCTAGTGCAGCCGGGA CCACACGCTATGTATGCCCGGGTGAGGCAGGTGGAGAGCAACCCCCTGTACCAGGGCACG GCCTCCAGCGCTGACGTGGCCCTGGTGGAGCTGGAGGCACCAGTGCCCTTCACCAATTAC ATCCTCCCCGTGTGCCTGCCTGACCCCTCGGTGATCTTTGAGACGGGCATGAACTGCTGG GTCACTGGCTGGGGCAGCCCCAGTGAGGAAGACCTCCTGCCGAACCGCGGATCCTGCAG AAACTCGCTGTGCCCATCATCGACACACCCAAGTGCAACCTGCTCTACAGCAAAGACACC GAGTTTGGCTACCAACCCAAAACCATCAAGAATGACATGCTGTGCGCCGGCTTCGAGGAG GGCAAGAAGGATGCCTGCAAGGGCGACTCGGGCGCCCCCTGGTGTGCCTCGTGGGTCAG TCGTGGCTGCAGGCGGGGTGATCAGCTGGGGTGAGGGCTGTGCCCGCCAGAACCGCCCA GGTGTCTACATCCGTGTCACCGCCCACCACAACTGGATCCATCGGATCATCCCCAAACTG CAGTTCCAGCCAGCGAGGTTGGGCGGCCAGAAGTCTAGACATCACCATCACCATCACTAG CGGCCGCTTCCCTTTAGTGAGGGTTAATGCTTCGAGCAGACATGATAAGATACATTGATG AGTTTGGACAAACCACAACTAGAATGCAGTGAAAAAAATGCTTTATTTGTGAAATTTGTG ATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTGAC

Amino Acid sequence of PFEK-PROTT-HIS fusion protein (SEQ.ID.NO.:9).

MDSKGSSQKSRLILLLVVSNLLLCQGVVSDYKDDDDVDAAALAAPFDDDDKIVGGYALEE GEWPWQVSIQRNGSHFCGGSLIAEQWVLTAAHCFRNTSETSLYQVLLGARQLVQPGPHAM YARVRQVESNPLYQGTASSADVALVELEAPVPFTNYILPVCLPDPSVIFETGMNCWVTGW GSPSEEDLLPEPRILQKLAVPIIDTPKCNLLYSKDTEFGYQPKTIKNDMLCAGFEEGKKD ACKGDSGGPLVCLVGQSWLQAGVISWGEGCARQNRPGVYIRVTAHHNWIHRIIPKLQFQP ARLGGQKSRHHHHHH



Protease: PFEK-Protease T-HIS

Protease Activity of Protease T

H-D-Pro-HHT-Arg-pNA
H-D-Lys(CBO)-Pro-Arg-pNA
H-DL-Val-Leu-Arg-pNA

